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Run
                                                        Database
                                                                                                                                                         Minimum DB seq length: 0
Maximum DB seq length: 2000000000
                                                                                                                                                                                                                                                                                                                                                      Title:
Perfect score:
                                                                                                                          Post-processing: Minimum Match 0%
                                                                                                                                                                                                            Total number of hits satisfying chosen parameters:
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Gapop 10.0 ,
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Listing first 45 summaries
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76
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                                                                                                                                                                                                                                              195891 seqs, 67900655 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GenCore version 4.5 Copyright (c) 1993 - 2000 Compugen Ltd
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pir2:*
pir3:*
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36.422 Million cell updates/sec
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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protein-tyrosine k	dna repair protein	IgG Fc binding pro	hypothetical prote	hypothetical prote	_	hypothetical prote	hypothetical prote	von Willebrand fac	diacylglycerol kin	diacylglycerol kin	hypothetical prote	hypothetical prote	hypothetical prote	hypothetical prote	insulin-like growt

ALIGNMENTS

A; Experimental source: clone C; Genetics: hypothetical protein C06A1.6 - Caenorhabditis elegans
C;Speckes: Caenorhabditis elegans
C;Date: 15-Oct-1999 #sequence_revision 15-Oct-1999 #text_change 15-Oct-1999
C;Accession: T18975 RESULT T18975 A;Gene: CESP:C06A1.6 A;Map position: 2 submitted to the EMBL Data Library, A; Reference number: Z19054 A; Accession: T18975 A; Status: preliminary; translated fi A; Map position: 3 C; Superfamily: fruit fly testis-specific protein C; Keywords: spermatogenesis; tandem repeat A;Cross-references: EMBL:X67703; NID:g11072; PIDN:CAA47938.1; PID:g11074 A;Note: sequence extracted from NCBI backbone (NCBIN:74217, NCBIP:74220) C;Genetics: Mech. Dev. 35, 143-151, 1991
A;Title: A cluster of four genes selectively expressed in the male germ line of Droso A;Reference number: A56565; MUID:92102953
A;Accession: S25773 testis-specific protein Mst84Db - fruit fly (Drosophila melanogaster)
C;Species: Drosophila melanogaster
C;Date: 26-Jul-1996 #sequence_revision 26-Jul-1996 #text_change 20-Aug-1999 A;Cross-references: EMBL:Z49886; PIDN:CAA90055.1; GSPDB:GN00020; CESP:C06A1.6 A;Experimental source: clone C06A1 A; Molecule type: DNA A; Residues: 1-152 <WIL> В Ş A;Gene: Mst84Db A;Cross-references: FlyBase:FBgn0004173 A; Molecule type: DNA A; Residues: 1-74 < KUH> R; McMurray, A. R;Kuhn, R.; Kuhn, C.; Boersch, D.; Glaetzer, K.H.; Schaefer, U.; Schaefer, M C; Accession: S25773; B56565 Query Match Best Local S Matches 6 Local Similarity 21.4 les 6; Conservative 12 CSPCGGPCGPCGPCGSCCSPCGSCC 39 preliminary; translated from GB/EMBL/DDBJ 100.0%; Score 76 21.4%; Pred. No. 22; Score 76; LE June 1995 Mismatches DB 2; Length 74; 0; Indels 0 Gaps 0

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RESULT
S25774
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A;Residues: 1-188 <NHA>
A;Cross-references: EMBL:U58760; NID:g1330384; PID:g1330389; PIDN:AAB00710.1; GSPDB:GN00A;Experimental source: strain Bristol N2; clone C27A2
                                                                                                                                                                                                                                                                                                                                                       submitted to the EMBL Data Library, May 1996
A; Description: The sequence of C. elegans cosmid C27A2
A; Reference number: Z18382
A; Accession: T15651
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C;Date: 20-Sep-1999 #sequence_revision
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A; Introns: 20/3; 90/2
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A; Residues: 1-164 <WIL>
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2; Mismatches
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RESULT
S59073
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C;Species: Scylla serrata (mud crab)

C;Date: 19-Feb-1984 #sequence_revision 19-Feb-1984

C;Accession: A03284

R;Lerch, K.; Ammer, D.; Olafson, R.W.

J. Biol. Chem. 257, 2420-2426, 1982

A;Title: Crab metallothionein. Primary structures

A;Reference number: A92363; MUID:82142340

A;Accession: A03284
                                                                                                                                                                                                                                  metallothionein isoform IIa - blue crab
C:Species: Callinectes sapidus (blue crab)
C:Date: 19-Mar-1997 #sequence_revision 19-Mar-1997 #text_change 07-May-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A;Cross references: EMBL:X67703; NID:g11072; PIDN:CAA47939.1; PID:g11075 A;NOte: the authors translated the codon TGC for residue 55 as Thr A;Note: sequence extracted from NCBI backbone (NCBIN:74217, NCBIP:74222) C;Genetics:
A;Gene: Mst84Dc
A;Cross references: FlyBase:FBgn0004174
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C;Superfamily: metallothionein
C;Keywords: metal binding
                                            A; Molecule type: protein A; Residues: 1-57 <BRO>
                                                                                            Biochem. J. 311, 617-622, 1995
A;Title: Primary structure and tissue-specific expression of blue crab (Callinectes A;Reference number: S59072; MUID:96033062
A;Accession: S59073
                                                                                                                                                                 R;Brouwer, M.; Enghild, J.; Hoexum-Brouwer, T.; Thogersen, Biochem. J. 311, 617-622, 1995
                                                                                                                                                                                                                 C; Accession: S59073
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A;Residues: 1-57 <LER>
C;Superfamily: metallothionein
C;Keywords: metal binding
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C; Superfamily: fruit fly testis-specific protein
C; Keywords: spermatogenesis; tandem repeat
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A;Title: A cluster of four genes selectively expressed in
A;Reference number: A56565; MUID:92102953
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A;Molecule type: DNA
A;Residues: 1-55 <KUH>
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A; Note: the C; Superfamil: C; Keywords: I
              metallothionein 1 - American lobster
C;Species: Homarus americanus (American lobster)
C;Date: 31-Jan-1992 #sequence_revision 31-Jan-1992 #text_change 12-Apr-1995
C;Accession: A37039
R;Brouwer, M.; Winge, D.R.; Gray, W.R.
J. Inorg. Biochem. 35, 289-303, 1989
A;Title: Structural and functional diversity of copper-metallothioneins from
                                                                                                                          RESULT
A37039
                                                                                                                                                                                                                                                                                                                               A; Molecule type: protein A; Residues: 1-58 <BRO> C; Superfamily: metalloth: C; Keywords: metal binding
                                                                                                                                                                                                                                                                                                                                                                                                      R;Brouwer, M.; Enghild, J.; Hoexum-Brouwer, T.; Thogersen, I.; Truncali, A. Biochem. J. 311, 617-622, 1995
A;Tille: Primary structure and tissue-specific expression of blue crab (Callinectes A;Reference number: S59072; MUID:96033062
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              C;Accession: A03283
R;Lerch, K.; Ammer, D.; Olafson, R.W.
J. Biol. Chem. 257, 2420-2426, 1982
A;Title: Crab metallothionein. Primary structures
A;Reference number: A92363; MUID:82142340
A;Accession: A03283
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       metallothionein isoform Ia - blue crab
C;Species: Callinectes sapidus (blue crab)
C;Date: 19-Mar-1997 #sequence_revision 19-Mar-1997 #text_change 07-May-1999
C;Accession: S59072
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A;Residues: 1-58 <LER>
A;Residues: 1-58 <LER>
C;Superfamily: metallothionein
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C:Species: Scylla serrata (mud crab)
C:Date: 29-Jul-1981 #sequence_revision
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Keywords: metal binding
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les 5; Conservative
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A; Residues: 1-63 < KUH>
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                Conservative
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A;Note: the sequence from Fig. 4 is inconsistent with that from Fig. 6 in h: C;Superfamily: metallothionein C;Keywords: metal binding; chelantion; metal-thiolate cluster C;Keywords: metal binding; chelantion; metal ions (Cys) #status predicted F;6,12,17,21,28/Binding site: transition metal ions (Cys) #status predicted F;31,38,46,47,54,56,57/Binding site: transition metal ions (Cys) #status predicted F;31,38,46,47,54,56,57/Binding site:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           testis-specific protein Mst84Da - fruit fly (Drosophila melanogaster)
C;Species: Drosophila melanogaster
C;Date: 26-Jul-1996 #seguence_revision 26-Jul-1996 #text_change 20-Aug-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   R;Pedersen, K.L.; Pedersen, S.N.; Hojrup, P.; Andersen, J.S.; Roepstorff, P.; Knudsen Biochem. J. 297, 60:614, 1994
A;Title: Purification and characterization of a cadmium-induced metallothionein from A;Reference number: S43367; MUID:94153337
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A;Residues: 1-58 <BRO>
C;Superfamily: metallothionein
                                                                                                          C;Superfamily: fruit fly testis-specific protein
C;Keywords: spermatogenesis; tandem repeat
                                                                                                                                                                                                                                                                    A;Cross-references: EMBL:X67703; NID:g11072; PIDN:CAA47937.1; PID:g11073
A;Note: sequence extracted from NCBI backbone (NCBIN:74219, NCBIP:74224)
                                                                                                                                                                                                                                                                                                                                                            A; Reference number: A56565; A; Accession: S25772
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A;Tille: A cluster of four genes selectively expressed in the male germ line
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C;Date: 19-Mar-1997 #sequence_revision 01-Feb-1999 #text_change 07-May-1999
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keratin I, feather - chicken
(;Species: Gallus gallus (chicken)
C;Date: 17-Dec-1982 #sequence_revision 12-Apr-1996 #text_change 22-Jun-1999
C;Accession: $06805; A02848
R;Presland, R.B.; Gregg, K.; Molloy, P.L.; Morris, C.P.; Crocker, L.A.; Rogers, G.E.
J. Mol. Biol. 209, 549-559, 1989
A;Title: Avian keratin genes. I. A molecular analysis of the structure and expression A;Reference number: $06805; MUID:90064515
A;Accession: $06805
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C;Species: Drosophila melanogaster
C;Date: 26-Jul-1996 #sequence_revision 26-Jul-1996 #text_change 20-Aug-1999
C;Accession: S25775; D56565
R;Kuhn, R.; Kuhn, C.; Boersch, D.; Glaetzer, K.H.; Schaefer, U.; Schaefer, M. Mech. Dev. 35, 143-151, 1991
A;Title: A cluster of four genes selectively expressed in the male germ line of Drosophi A; Reference number: A56565; MUID:92102953
A;Accession: S25775
A;More in the male germ line of Drosophi A;Accession: S25775
A;More in the male germ line of Drosophi A;Accession: S25775
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A;Title: Organisation of feather keratin genes in the chick genome. A;Reference number: A02848; MUID:83064534
A;Accession: A02848
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A;Cross-references: FlyBase:FBgn0004175
A;Map position: 3
C;Superfamily: fruit fly testis-specific protein
C;Keywords: spermatogenesis; tandem repeat
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A; Residues: 2-98 < MOL>
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A; Residues: 1-98 < PRE>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A; Cross-references: GB:J00847; NID:g212229; PIDN:AAA48930.1; PID:g212230
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     밁
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A; Status: not compared with conceptual translation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Note: sequence extracted C;Genetics:
                                                                       Query Match
Best Local Similarity
Matches 5; Conserv
                                                                                                                                                                                                                                                                                                                          Superfamily:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     13 CGPCCGPCCGPCGPCGPCGPC 39
                                                                                                                                                                                                                                                                                                                                                                                                                                                a genomic clone containing five feather keratin genes was isolated and the cent.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        14 GPCCGPCGGCGPCGGCGPCC 40
::|:::|::::|::::|:::|
                                                                                                                                                                                                                                                                        ly: feather keratin epidermis; feather; fibrous protein; horn; integument
                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                     88.2%;
18.5%;
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18.5%; Pred. No.
                                                                               22;
                                                                                                                     Score 67; DB 1;
Pred. No. 1.2e+02;
                                                                           Mismatches
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96;
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                                                                                                                                                       Length 98
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                                                                               Indels
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                                                                                                                                                                  A;Cross-references: GB:X00316; NID:g63550; PIDN:CAA25085.1; PID:g63551 C;Superfamily: feather keratin C;Keywords: epidermis; feather; fibrous protein; horn; integument
                                                                                                                                                                                                                            A;Title: A comparison of genomic coding sequences for feather and A;Reference number: A90988; MUID:84158528
A;Accession: A02849
A;Molecule type: DNA
A;Residues: 1-98 <GRE>
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                                                                                                                                                                                                                                                                                                                           R;Gregg, K.; Wilton, S.D.; Parry, D.A.D.; Rogers, G.E. EMBO J. 3, 175-178, 1984
                                                                                                                                                                                                                                                                                                                                                                                   C;Species: Gallus gallus (chicken)
C;Date: 17-Mar-1987 #sequence_revision 17-Mar-1987 #text_change 22-Jun-1999
                                                                                                                                                                                                                                                                                                                                                                   C; Accession: A02849
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Best Local
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MSCYDLCRPCGPTPLANSCNEPCVRQC 27
                                  MSCEDLCRPCGPTPLANSCNEPCVRQC
                                                                                                                                                                                                                                                                                                                                                                                                                         feather - chicken
                                                                        Similarity 18.9
5; Conservative
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18.5%; Pred. No. 1
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